. Tambel

PAGE: 1

RAW SEQUENCE LISTING PATENT APPLICATION US/08/477,989B

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This Raw Listing contains the General Information Section and up to the first 5 pages.

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			× 70.
1		SEQUENCE LISTING	as E.
2		7 7 6 11	
3	(1)	General Information:	1 /2
<u>4</u> 5		(i) APPLICANT: Bazin, Herv	
6		Latinne, Dominique	
7		Kaplan, Ruth Kieber-Emmons, Thom	
8		Postema, Christina	ds T
9			<u> </u>
10		White-Scharf, Mary (ii) TITLE OF INVENTION: LO-CD2a Antibo	de and Hann
11		(ii) TITLE OF INVENTION: LO-CD2a Antibo Thereof for In	
12		T-Cell Activat	
13		Proliferation	1011 and
14		(iii) NUMBER OF SEQUENCES: 96	
15		(iv) CORRESPONDENCE ADDRESS:	
16		(A) ADDRESSEE: Carella, Byrne, Bai.	n Gilfillan
17		Cecchi, Stewart & O.	
18		(B) STREET: 6 Becker Farm Road	IPCeill
19		(C) CITY: Roseland	
20		(D) STATE: New Jersey	
21		(E) COUNTRY: U.S.A.	
22		(F) ZIP: 07068	RECT
23		(1) 211. 0/000	RECEIVED
24		(v) COMPUTER READABLE FORM:	
25		(A) MEDIUM TYPE: 3.5 inch diskette	APR 0 4 1999
26		(B) COMPUTER: IBM PS/2	71 K 0 4 1900
27		(C) OPERATING SYSTEM: MS-DOS	1009
28		(D) SOFTWARE: WordPerfect 5.1	•
29		,	
30		(vi) CURRENT APPLICATION DATA:	
31		(A) APPLICATION NUMBER: 08/477,989	
32		(B) FILING DATE: 07-JUN-1995	
33		(C) CLASSIFICATION:	
34		, ,	
35		(vii) PRIOR APPLICATION DATA:	
36		(A) APPLICATION NUMBER: 08/407,009	
37		(B) FILING DATE: 29-MAR-1995	
38		(A) APPLICATION NUMBER: 08/119,032	_
39		(B) FILING DATE: 09-SEP-1993	a g
40		(A) APPLICATION NUMBER: 08/027,008	
41		(B) FILING DATE: 05-MAR-1993	S O (1)
42			
43		(viii) ATTORNEY/AGENT INFORMATION:	[4] 1 28
44		(A) NAME: Olstein, Elliot M.	APR 7 - 19
45		(B) REGISTRATION NUMBER: 24,025	
46		(C) REFERENCE/DOCKET NUMBER: 61750-	147 🔐 🛱 📆
			APR Corrections
			රි
			-

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47		
48	(ix) TELECOMMUNICATION INFORMATION:	
49	(A) TELEPHONE: 201-994-1700	
50	(B) TELEFAX: 201-994-1744	
51		
52	(2) INFORMATION FOR SEQ ID NO: 1:	
53	(i) SEQUENCE CHARACTERISTICS:	
54	(A) LENGTH: 32 bases	
55	(B) TYPE: nucleic acid	
56	(C) STRANDEDNESS: single	
57	(D) TOPOLOGY: linear	
58	(ii) MOLECULE TYPE: oligonucleotide	
59		
60	(ix) FEATURE:	
61	(A) NAME/KEY: PCR primer	
62	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:	
63	AACCCGGGGA CATTCAGCTG ACCCAGTCTC AA 32	2
64		
65	(2) INFORMATION FOR SEQ ID NO: 2:	
66	(i) SEQUENCE CHARACTERISTICS:	
67	(A) LENGTH: 29 bases	
68	(B) TYPE: nucleic acid	
69	(C) STRANDEDNESS: single	
70	(D) TOPOLOGY: linear	
71	(ii) MOLECULE TYPE: oligonucleotide	
72		
73	(ix) FEATURE:	
74	(A) NAME/KEY: PCR primer	
75	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:	
75 76		9
75 76 77	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC 29	9
75 76 77 78	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC 29 (2) INFORMATION FOR SEQ ID NO: 3:	9
75 76 77 78 79	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC 29 (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS:	9
75 76 77 78 79 80	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases	9
75 76 77 78 79 80 81	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid	•
75 76 77 78 79 80 81 82	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single	•
75 76 77 78 79 80 81 82 83	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC 29 (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	9
75 76 77 78 79 80 81 82 83	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single	9
75 76 77 78 79 80 81 82 83 84 85	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide	9
75 76 77 78 79 80 81 82 83 84 85	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide (ix) FEATURE:	9
75 76 77 78 79 80 81 82 83 84 85 86	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC 29 (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide (ix) FEATURE: (A) NAME/KEY: PCR primer	9
75 76 77 78 79 80 81 82 83 84 85 86 87	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide (ix) FEATURE: (A) NAME/KEY: PCR primer (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:	
75 76 77 78 79 80 81 82 83 84 85 86 87 88	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC 29 (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide (ix) FEATURE: (A) NAME/KEY: PCR primer	
75 76 77 78 79 80 81 82 83 84 85 86 87 88 89	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide (ix) FEATURE: (A) NAME/KEY: PCR primer (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3: AACCCGGGGA GGTCCAGCTG CAGCAGTCTG G 31	
75 76 77 78 79 80 81 82 83 84 85 86 87 88 90 91	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC 29 (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide (ix) FEATURE: (A) NAME/KEY: PCR primer (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3: AACCCGGGGA GGTCCAGCTG CAGCAGTCTG G 31 (2) INFORMATION FOR SEQ ID NO: 4:	
75 76 77 78 79 80 81 82 83 84 85 86 87 88 90 91 92	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide (ix) FEATURE: (A) NAME/KEY: PCR primer (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3: AACCCGGGGA GGTCCAGCTG CAGCAGTCTG G (2) INFORMATION FOR SEQ ID NO: 4: (i) SEQUENCE CHARACTERISTICS:	
75 76 77 78 79 80 81 82 83 84 85 86 87 88 90 91 92 93	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide (ix) FEATURE: (A) NAME/KEY: PCR primer (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3: AACCCGGGGGA GGTCCAGCTG CAGCAGTCTG G 31 (2) INFORMATION FOR SEQ ID NO: 4: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 bases	
75 76 77 78 79 80 81 82 83 84 85 86 87 88 90 91 92 93	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide (ix) FEATURE: (A) NAME/KEY: PCR primer (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3: AACCCGGGGGA GGTCCAGCTG CAGCAGTCTG G 31 (2) INFORMATION FOR SEQ ID NO: 4: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 bases (B) TYPE: nucleic acid	
75 76 77 78 79 80 81 82 83 84 85 86 87 88 90 91 92 93 94 95	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide (ix) FEATURE: (A) NAME/KEY: PCR primer (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3: AACCCGGGGA GGTCCAGCTG CAGCAGTCTG G 31 (2) INFORMATION FOR SEQ ID NO: 4: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single	
75 76 77 78 79 80 81 82 83 84 85 86 87 88 90 91 92 93 94 95 96	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide (ix) FEATURE: (A) NAME/KEY: PCR primer (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3: AACCCGGGGA GGTCCAGCTG CAGCAGTCTG G 31 (2) INFORMATION FOR SEQ ID NO: 4: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
75 76 77 78 79 80 81 82 83 84 85 86 87 88 90 91 92 93 94 95 96 97	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide (ix) FEATURE: (A) NAME/KEY: PCR primer (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3: AACCCGGGGA GGTCCAGCTG CAGCAGTCTG G 31 (2) INFORMATION FOR SEQ ID NO: 4: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single	
75 76 77 78 79 80 81 82 83 84 85 86 87 88 90 91 92 93 94 95 96	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2: CAGTCGACTA CAGTTGGTGC AGCATCAGC (2) INFORMATION FOR SEQ ID NO: 3: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 31 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: oligonucleotide (ix) FEATURE: (A) NAME/KEY: PCR primer (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3: AACCCGGGGA GGTCCAGCTG CAGCAGTCTG G 31 (2) INFORMATION FOR SEQ ID NO: 4: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 bases (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	

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		INFULSEL: 32/2
100	(A) NAME/KEY: PCR primer	
101	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:	0.7
102	AAGTCGACCC AGTGGATAGA CCGATGG	27
103	/A) THEODINETON FOR GEO TO NO. F.	
104 105	(2) INFORMATION FOR SEQ ID NO: 5:(i) SEQUENCE CHARACTERISTICS:	
105	(A) LENGTH: 22 bases	
105	(B) TYPE: nucleic acid	
107	(C) STRANDEDNESS: single	
100	(D) TOPOLOGY: linear	
110	(ii) MOLECULE TYPE: oligonucleotide	
111	(11) Holdon 1111 Calgoria	
112	(ix) FEATURE:	
113	(A) NAME/KEY: PCR primer	
114	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:	
115	CCGCAAGCTT CATGGGATGG AG	22
116		
117	(2) INFORMATION FOR SEQ ID NO: 6:	
118	(i) SEQUENCE CHARACTERISTICS:	
119	(A) LENGTH: 27 bases	•
120	(B) TYPE: nucleic acid	
121	(C) STRANDEDNESS: single	
122	(D) TOPOLOGY: linear	
123	(ii) MOLECULE TYPE: oligonucleotide	
124	(in) DELEGIOE.	
125	(ix) FEATURE:	
126 127	(A) NAME/KEY: PCR primer (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:	
127	GCTGCTTGGG GACTGGGTCA GCTGGAT	27
129	GCIGCIIGGG GACIGGGICA GCIGGAI	2,
130	(2) INFORMATION FOR SEQ ID NO: 7:	
131	(i) SEQUENCE CHARACTERISTICS:	
132	(A) LENGTH: 21 bases	
133	(B) TYPE: nucleic acid	
134	(C) STRANDEDNESS: single	
135	(D) TOPOLOGY: linear	
136	(ii) MOLECULE TYPE: oligonucleotide	
137		
138	(ix) FEATURE:	
139	(A) NAME/KEY: PCR primer	
140	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:	
141	ATTCAGCTGA CCCAGTCTCC A	21
142	(2) THEODYSETON FOR GEO TO NO. 9.	
143	(2) INFORMATION FOR SEQ ID NO: 8: (i) SEQUENCE CHARACTERISTICS:	
144 145	(1) SEQUENCE CHARACTERISTICS: (A) LENGTH: 58 bases	
145	(A) LENGTH: 50 bases (B) TYPE: nucleic acid	
147	(C) STRANDEDNESS: single	
148	(D) TOPOLOGY: linear	
149	(ii) MOLECULE TYPE: oligonucleotide	
150	(,	
151	(ix) FEATURE:	
152	(A) NAME/KEY: PCR primer	
	• •	

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153	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
154	GATCGGATCC ACCTGAGGAA GCAAAGTTTA AATTCTACTC ACGTTTCAGT TCCAGCTT58
155	(A) THROPHIMTON FOR GEO TO NO. A.
156	(2) INFORMATION FOR SEQ ID NO: 9:
157 158	(i) SEQUENCE CHARACTERISTICS:
	(A) LENGTH: 31 bases
159	(B) TYPE: nucleic acid
160	(C) STRANDEDNESS: single
161	(D) TOPOLOGY: linear
162 163	(ii) MOLECULE TYPE: oligonucleotide
164	(ix) FEATURE:
165	(A) NAME/KEY: PCR primer
166	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
167	TCTCCTGCAG TGGGACCTCG GAGTGGACAC C 31
168	TOTOCOGCAG TOGGACCTCG GAGTGGACAC C 51
169	(2) INFORMATION FOR SEQ ID NO: 10:
170	(i) SEQUENCE CHARACTERISTICS:
171	(A) LENGTH: 21 bases
172	(B) TYPE: nucleic acid
173	(C) STRANDEDNESS: single
174	(D) TOPOLOGY: linear
175	(ii) MOLECULE TYPE: oligonucleotide
176	(11) 1101110011 11111 0119011110
177	(ix) FEATURE:
178	(A) NAME/KEY: PCR primer
179	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
180	GAGGTCCAGC TGCAGCAGTC T 21
181	
182	(2) INFORMATION FOR SEQ ID NO: 11:
183	(i) SEQUENCE CHARACTERISTICS:
184	(A) LENGTH: 24 bases
185	(B) TYPE: nucleic acid
186	(C) STRANDEDNESS: single
187	(D) TOPOLOGY: linear
188	(ii) MOLECULE TYPE: oligonucleotide
189	
190	(ix) FEATURE:
191	(A) NAME/KEY: PCR primer
192	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
193	CGATGTATCA GCTGTCAGTG TGGC 24
194	
195	(2) INFORMATION FOR SEQ ID NO: 12:
196	(i) SEQUENCE CHARACTERISTICS:
197	(A) LENGTH: 24 bases
198	(B) TYPE: nucleic acid
199	(C) STRANDEDNESS: single
200	(D) TOPOLOGY: linear
201	(ii) MOLECULE TYPE: oligonucleotide
202	/irr\ BEADUDE.
203	(ix) FEATURE:
204	(A) NAME/KEY: PCR primer
205	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:

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(A) NAME/KEY: PCR primer

TTGGATCCGC GGCCGCGTCG ACTACAGTTG GTGCAGCATC AGC

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:

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